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Amendments to the Claims:

This listing of claims will replace all prior versions, and listings of claims in the application:

Listing of Claims:

- 1. (Amended) A method of determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide or a thioambenyl ethionamide, thiacetazone or thiocarlide, said method comprising detecting a mutation in an EtaA gene (SEQ ID NO:1) in said bacterium, which mutated gene encodes an amino acid sequence which differs from that of SEQ ID NO:2 by
- (a) a frameshift mutation selected from the group consisting of: a deletion at position 65, an addition at position 567, and an addition at position 811, or
- (b) a single nucleotide polymorphism which causes an amino acid substitution selected from the group consisting of: G43C, P51L, D58A, Y84D, T186K, T342K, and A381P, wherein detection of the mutation is indicative of decreased ability to oxidize a thioamide or a thiocarbonyl ethionamide, thiacetazone or thiocarlide.
 - 2. (Canceled)
- 3. (Original) The method of claim 1, wherein the mutation is a single nucleotide polymorphism which causes an amino acid substitution in an amino acid sequence encoded by said EtaA gene compared to an amino acid sequence of SEQ ID NO:2.
 - 4. (Canceled)
- 5. (Original) A method of claim 1 wherein the mutation is detected by

 (a) amplifying the EtaA gene, or a portion thereof containing the mutation, with a set of primers to provide an amplified product,
 - (b) sequencing the amplified product to obtain a sequence, and

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(c) comparing the sequence of the amplified product with the sequence of a wild-type EtaA gene (SEQ ID NO:1) or portion thereof,

wherein a difference between the sequence of the amplified product and the sequence of the wild-type EtaA gene or portion thereof indicates the presence of a mutation.

- 6-7. Canceled.
- 8. (Original) A method of claim 5, wherein said amplification is by polymerase chain reaction.
- 9. (Original) A method of claim 1, wherein said mutation is detected by hybridizing DNA from said bacterium to a test nucleic acid under stringent conditions.
- 10. (Original) A method of claim 9, wherein either said DNA from said bacterium or said test nucleic acid is immobilized on a solid support.
 - 11. (Original) A method of claim 1, wherein said mutation is detected by
 - (a) subjecting said EtaA gene to digestion by restriction enzymes,
- (b) separating the resulting restriction products to form a pattern of restriction fragment lengths, and
- (c) comparing the pattern of restriction fragment lengths to a pattern of restriction fragment lengths formed by subjecting a known EtaA gene to the same restriction enzymes.
 - 12. (Canceled)
- 13. (Withdrawn) A method of claim 1, wherein said mutation is detected by specifically binding an antibody to a mutated product of the EtaA gene, wherein the specific binding of the antibody to the mutated gene product is indicative of a mutation which inhibits the ability of the bacterium to oxidize a thioamide.

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- 14. (Withdrawn) A method of claim 13, wherein said gene product is in, or is isolated from, sputum.
- 15. (Withdrawn) A method of claim 13, wherein detection of said specific binding of said antibody and said mutated gene product is by ELISA.
 - Canceled.
 - . 17. (Withdrawn) A method of claim 1, wherein said mutation is detected by
 - (a) culturing said bacterium in the presence of ethionamide; and
- (b) testing for the presence or absence of (2-ethyl-pyridin-4-yl)methanol, wherein the absence of (2-ethyl-pyridin-4-yl)methanol indicates that the bacterium has a mutation which is indicative of decreased ability to oxidize a thioamide.
- (2-ethyl-pyridin-4-yl)methanol is tested by subjecting a medium in which the bacterium is cultured, or the bacterium, to analysis by thin-layer chromatography, high pressure liquid chromatography, or mass spectrometry.
- (Withdrawn) A method of claim 17, wherein the ethionamide of step (a) is radioactively labeled.
- 20. (Withdrawn) A method of claim 17, wherein the (2-ethyl-pyridin-4-yl)methanol is radioactively labeled.
- 21. (Currently amended) A method of screening an individual for a Mycobacterium tuberculosis bacterium resistant to treatment by a thioamide or a thiocarbonyl drug ethionamide, thiacetazone or thiocarlide, comprising
- (a) obtaining a biological sample containing said bacterium from said individual, and
 - (b) detecting a mutation in an EtaA gene (SEQ ID NO:1) in said bacterium, which

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mutated gene encodes an amino acid sequence which differs from that of SEQ ID NO;2, wherein said mutation in said EtaA gene is selected from the group consisting of

(i) a frameshift mutation consisting of a deletion at position 65, an addition at position 567, or an addition at position 811, and

(ii) a single nucleotide polymorphism which causes an arnino acid substitution selected from the group consisting of: G43C, P51L, D58A, Y84D, T186K, T342K, and A381P, wherein detection of the mutation is indicative said bacterium is resistant to treatment by a thioamide or a thiocarbonyl drug ethionamide, thiacetazone or thiocarlide.

- (Original) A method of claim 21, wherein the mutation is detected by

 (a) amplifying the EtaA gene with a set of primers to provide an amplified product,
 - (b) sequencing the amplified product to obtain a sequence, and
- (c) comparing the sequence of the amplified product with the sequence of a wild-type EtaA gene (SEQ ID NO:1).

wherein a difference between the sequence of the amplified product and the sequence of the wild-type EtaA gene indicates the presence of a mutation.

23-24. Canceled.

- 25. (Previously presented) A kit for determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide or a thiocarbonyl ethionamide, thiacetazone or thiocarlide, the kit comprising:
 - (a) a container, and
- (b) primers for specifically amplifying an EtaA gene of said bacterium or a portion of said EtaA gene containing a mutation affecting the ability of the bacterium to exidize a thioamide selected from the group consisting of (i) a frameshift mutation consisting of a deletion at position 65, an addition at position 567, or an addition at position 811, and

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- (ii) a single nucleotide polymorphism which causes an amino acid substitution selected from the group consisting of: G43C, P51L, D58A, Y84D, T186K, T342K, and A381P.
 - 26-27. Canceled.
- 28. (Original) A kit of claim 25, further comprising a mutated EtaA gene for use as a positive control.
 - 29. (Canceled)
- 30. (Withdrawn) A kit for determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide, the kit comprising:
 - (a) a container, and
 - (b) (2-ethyl-pyridin-4-yl)methanol.
- 31. (Withdrawn) A kit for determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide, the kit comprising:
 - (a) a container, and
 - (b) radiolabeled ethioamide.
- 32. (Withdrawn) A kit for determining the ability of a *Mycobacterium* tuberculosis bacterium to oxidize a thioamide or thiocarbonyl, the kit comprising:
 - (a) a container, and
- (b) an antibody which specifically binds to a product of a EtaA gene selected from the group consisting of a wild-type EtaA gene (SEQ ID NO:1) and a mutated EtaA gene.
- 33. (Withdrawn) A kit for determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide, the kit comprising:
 - (a) a container, and
 - (b) an antibody which specifically binds to (2-ethyl-pyridin-4-yl)methanol.

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- 34. (Currently amended) A method of determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide or a thiocarbonyl selected from the group consisting of ethionamide, thiacetazone and thiocarlide, said method comprising detecting a mutation in an EtaA gene (SEQ ID NO:1) in said bacterium, which mutated gene encodes an amino acid sequence which differs from that of SEQ ID NO:2, wherein said mutation is selected from the group consisting of
- (a) a frameshift mutation consisting of a deletion at position 65, an addition at position 567, or an addition at position 811, and
- (b) a single nucleotide polymorphism which causes an amino acid substitution selected from the group consisting of: G43C, P51L, D58A, Y84D, T186K, T342K, and A381P.

wherein detection of the mutation is indicative of decreased ability to oxidize ethionamide, thiacetazone or thiocarlide.

- 35. (Previously presented) The method of claim 34, wherein the mutation is a frameshift mutation selected from the group consisting of: a deletion at position 65, an addition at position 567, and an addition at position 811.
 - 36. (Canceled)
- 37. (Currently amended) The method of claim 36 34, wherein the single nucleotide polymorphism causes an amino acid substitution selected from the group consisting of: G43C, P5 IL, D58A, Y84D, T186K, T342K, and A381P.
- 38. (Previously presented) A method of claim 34 wherein the mutation is detected by
- (a) amplifying the EtaA gene, or a portion thereof containing the mutation, with a set of primers to provide an amplified product,
 - (b) sequencing the amplified product to obtain a sequence, and

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(c) comparing the sequence of the amplified product with the sequence of a wild-type EtaA gene (SEQ ID NO:1) or portion thereof,

wherein a difference between the sequence of the amplified product and the sequence of the wild-type EtaA gene or portion thereof indicates the presence of a mutation.

- 39. (Previously presented) A method of claim 38, wherein said amplification is by polymerase chain reaction.
- 40. (Previously presented) A method of claim 34, wherein said mutation is detected by hybridizing DNA from said bacterium to a test nucleic acid under stringent conditions.
- 41. (Previously presented) A method of claim 40, wherein either said DNA from said bacterium or said test nucleic acid is immobilized on a solid support.
- 42. (Previously presented) A method of claim 34, wherein said mutation is detected by
 - (a) subjecting said EtaA gene to digestion by restriction enzymes,
- (b) separating the resulting restriction products to form a pattern of restriction fragment lengths, and
- (c) comparing the pattern of restriction fragment lengths to a pattern of restriction fragment lengths formed by subjecting a known EtaA gene to the same restriction enzymes.
 - 43. (Canceled)
- 44. (Currently amended) A method of screening an individual for a Mycobacterium tuberculosis bacterium resistant to treatment by a thioamide or a thiocarbonyl drug, selected from the group consisting of ethionamide, thiacetazone and thiocarlide, comprising

product,

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- (a) obtaining a biological sample containing said bacterium from said individual,
- (b) detecting a mutation in an EtaA gene (SEQ ID NO:1) in said bacterium, wherein said mutation is selected from the group consisting of (a) a frameshift mutation consisting of a deletion at position 65, an addition at position 567, and an addition at position 811, and (b) a single nucleotide polymorphism which causes an amino acid substitution selected from the group consisting of: G43C, P51L, D58A, Y84D, T186K, T342K, and A381P, wherein detection of the mutation is indicative said bacterium is resistant to treatment by ethionamide, thiacetazone or thiocarlide.
- 45. (Previously presented) A method of claim 44, wherein the mutation is detected by
 - (a) amplifying the EtaA gene with a set of primers to provide an amplified
 - (b) sequencing the amplified product to obtain a sequence, and
- (c) comparing the sequence of the amplified product with the sequence of a wild-type EtaA gene (SEQ ID NO:1),

wherein a difference between the sequence of the amplified product and the sequence of the wild-type EtaA gene indicates the presence of a mutation.

- 46. (Previously presented) A kit for determining the ability of a Mycobacterium tuberculosis bacterium to oxidize a thioamide or a thiocarbonyl selected from the group consisting of ethionamide, thiacetazone and thiocarlide, the kit comprising:
 - (a) a container, and
- (b) primers <u>specific</u> for amplifying an EtaA gene of said bacterium or a portion of said EtaA gene containing a mutation affecting the ability of the bacterium to oxidize ethionamide, thiacetazone or thiocarlide